

AMENDMENTS TO THE SPECIFICATION

Please replace the paragraph beginning at page 2, line 26, with the following amended paragraph:

—Furthermore, the present inventors searched cDNA sequences of Helix Research Institute (helix clones; Japanese Patent Application No. Hei 11-248036; Japanese Patent Application No. 2000-118776) for homologues using the above-mentioned sequence, “Hs128045_12F08con”, obtained from Unigene Cluster as a query. These helix clones are highly expected to have the full length sequence, which are obtained by the combined use of: [1] preparation of cDNA library, which comprises cDNA having a full-length sequence at a high rate, by the oligo-capping method; and [2] evaluation system for the cDNA to determine whether it contains the full-length sequence based on the 5' end sequence (the selection is achieved based on the evaluation using ATGpr after eliminating non-full length clones as compared with an EST). The results of homology search showed that the query clone was identical to the helix clone ~~“C-T2RP3001495”~~ “C-NT2RP3001495”. In addition, it was also revealed that the query clone is identical to the gene for Hs.519 Human oxidoreductase (HHCMA56) of Unigene. However, the sequence of HHCMA56 contains reading mistakes of nucleotides, and thus it has been deposited as a gene encoding a protein consisting of 371 amino acids which is entirely different from the protein of “C-NT2RP3001495”. Thus, it can be stated that “C-NT2RP3001495” is a novel protein found for the first time by the present inventors. The “C-NT2RP3001495” is a protein consisting of 414 amino acids, which has two WW domain sequences.—